

**In the Claims**

This listing of claims will replace all prior versions, and listings, of claims in the application:

Claims 1-3 (Cancelled)

4. (Previously presented) A transformed plant having a nucleic acid molecule which comprises:
  - (a) an exogenous promoter region which functions in a plant cell to cause the production of an mRNA molecule;
  - (b) a structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 and complement thereof; and
  - (c) a 3' non-translated sequence that functions in said plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule.
5. (Previously presented) The transformed plant according to claim 4, wherein said structural nucleic acid molecule is a complement of the nucleic acid sequence of SEQ ID NO: 1.
6. (Original) The transformed plant according to claim 5, wherein said plant is soybean or maize.
7. (Original) The transformed plant according to claim 5, wherein said plant is soybean.

8. (Previously presented) A method for determining a level or pattern in a plant cell or plant tissue of a protein in a plant comprising:
  - (a) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, said marker nucleic acid molecule selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having the nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 or complement thereof, with a complementary nucleic acid molecule obtained from said plant cell or plant tissue, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue permits the detection of an mRNA for said protein;
  - (b) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue; and
  - (c) detecting the level or pattern of said complementary nucleic acid, wherein the detection of said complementary nucleic acid is predictive of the level or pattern of said protein.
9. (Original) The method of claim 8, wherein said level or pattern is detected by *in situ* hybridization.
10. (Original) The method of claim 8, wherein said level or pattern is detected by tissue printing.
11. (Original) The method of claim 8, wherein said plant is maize or soybean.

12. (Original) The method of claim 11, wherein said plant is soybean.

Claims 13-17 (Cancelled)